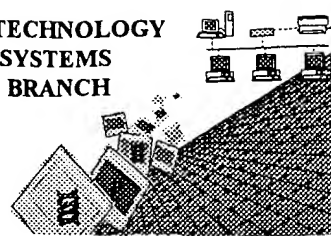


1653

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TECH CENTER 1600/2900

BIOTECHNOLOGY  
SYSTEMS  
BRANCH

## RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/105,117I  
Source: 1620  
Date Processed by STIC: 8/12/2002

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
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FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: [patin21help@uspto.gov](mailto:patin21help@uspto.gov) or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: [patin3help@uspto.gov](mailto:patin3help@uspto.gov) or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
3. Hand Carry directly to:  
U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7<sup>th</sup> Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202  
Or  
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 01/29/2002



1600

## RAW SEQUENCE LISTING

DATE: 08/12/2002

PATENT APPLICATION: US/09/105,117I

TIME: 14:05:45

Input Set : A:\Sequencelisting\_FZJ 9910 PCT\_US.txt

Output Set: N:\CRF3\08122002\I105117I.raw

**Does Not Comply  
Corrected Diskette Needed**

3 <110> APPLICANT: Forschungszentrum Juelich GmbH  
 5 <120> TITLE OF INVENTION: Process for the microbial production of amino acids by  
 6 boosted activity of export carriers  
 8 <130> FILE REFERENCE: 1  
 C--> 10 <140> CURRENT APPLICATION NUMBER: US/09/105,117I  
 C--> 11 <141> CURRENT FILING DATE: 1998-06-17  
 E--> 13 <160> NUMBER OF SEQ ID NOS: 25 (see p.4)  
 15 <170> SOFTWARE: PatentIn Ver. 2.0

## ERRORED SEQUENCES

E--> 206 <210> SEQ ID NO: DNA (complement to <210> 1)  
 207 <213> ORGANISM: Corynebacterium glutamicum  
 W--> 208 <220> FEATURE:  
 209 <221> NAME/KEY: unsure  
 210 <222> LOCATION: CDS (2)..(652)  
 211 <223> OTHER INFORMATION: orf3  
 W--> 212 <220> FEATURE:  
 213 <221> NAME/KEY: gene  
 214 <222> LOCATION: CDS (1421)..(2293)  
 215 <223> OTHER INFORMATION: LysG  
 W--> 217 <212> TYPE:  
 E--> 217 <400> SEQUENCE: 3

*see  
p.6 for explanation*

218 a gat act cct ttg gaa gaa acc atg tac gca ttg cgt gac att gtt gcg 49  
 219 Asp Thr Pro Leu Glu Glu Thr Met Tyr Ala Leu Arg Asp Ile Val Ala  
 220 1 5 10 15  
 222 tct gga aag gct ctt tac gtg ggt att tct tcc tac ggt cca gag ctc 97  
 223 Ser Gly Lys Ala Leu Tyr Val Gly Ile Ser Ser Tyr Gly Pro Glu Leu  
 224 20 25 30  
 226 aca gcg gag gcg gct gag ttc atg gcg gag gag ggc tgc ccg ctt ctg 145  
 227 Thr Ala Glu Ala Ala Glu Phe Met Ala Glu Glu Gly Cys Pro Leu Leu  
 228 35 40 45  
 230 att cat cag cca agc tat tcc atc att aat cgt tgg gtg gag gaa ccg 193  
 231 Ile His Gln Pro Ser Tyr Ser Ile Ile Asn Arg Trp Val Glu Glu Pro  
 232 50 55 60  
 234 ggc gat gac ggt gag aac ttg ttg cag tca gct gcc aac aat ggt ctt 241  
 235 Gly Asp Asp Gly Glu Asn Leu Leu Gln Ser Ala Ala Asn Asn Gly Leu  
 236 65 70 75 80  
 238 ggc gtc att gct ttc tca cca ctt gcg cag ggc ctg ctc acg gac aaa 289  
 239 Gly Val Ile Ala Phe Ser Pro Leu Ala Gln Gly Leu Leu Thr Asp Lys  
 240 85 90 95  
 242 tat ctc gat gga att cca gag ggt tcc cgc gcc agc cag ggt aag tcc 337

## RAW SEQUENCE LISTING

DATE: 08/12/2002

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TIME: 14:05:45

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Output Set: N:\CRF3\08122002\I105117I.raw

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243 Tyr Leu Asp Gly Ile Pro Glu Gly Ser Arg Ala Ser Gln Gly Lys Ser
244      100      105      110
246 ctg tct gag ggc atg ttg aac gtg aac aat att gat atg gtc cgc aag 385
247 Leu Ser Glu Gly Met Leu Asn Val Asn Asn Ile Asp Met Val Arg Lys
248      115      120      125
250 ctc aat gac atc gcc cag gaa cgc ggg cag tca ctt gcg cag atg gcg 433
251 Leu Asn Asp Ile Ala Gln Glu Arg Gly Gln Ser Leu Ala Gln Met Ala
252      130      135      140
254 ctt gca tgg gtg ctg cgc gag caa gga gag tac ggc gcg gat acc gtg 481
255 Leu Ala Trp Val Leu Arg Glu Gln Gly Glu Tyr Gly Ala Asp Thr Val
256 145      150      155      160
258 acc agt gca ttg att ggt gct tcg tca gtt gag cag ctg gac aac agc 529
259 Thr Ser Ala Leu Ile Gly Ala Ser Ser Val Glu Gln Leu Asp Asn Ser
260      165      170      175
262 ctt gat tca ctc aac aac ttg gag ttt tct gac gcc gag ttg gag gcg 577
263 Leu Asp Ser Leu Asn Asn Leu Glu Phe Ser Asp Ala Glu Leu Glu Ala
264      180      185      190
266 atc gat gag att tcc cac gac gcc ggc atc aac att tgg gcg aag gcc 625
267 Ile Asp Glu Ile Ser His Asp Ala Gly Ile Asn Ile Trp Ala Lys Ala
268      195      200      205
270 acc gat tcc aaa acc cgc gaa aac taa cccatcaaca tcagtttgat 672
271 Thr Asp Ser Lys Thr Arg Glu Asn
272      210      215
274 ggccaatgcg gtcatacaca ctgccacgac gacgttgatc cagcgccaca ccttggggct 732
276 ggacagcggg cgtgacaatg ctgctgcgcc gaaacccacc agcgggaacc agatcaggct 792
278 tgccgcgaac gcgccagcgg cgaaaatcca ccgtccggtg tcgccgtatt gcgcgccgac 852
280 gccgccgata aacacaaaac cgtccaaata cgcattcggg ttcaaccagg tcagcacgat 912
282 tgccatcaac atgggcttta cccaaacccg ctgcttatcg acgctcacct ccaccgcac 972
284 ccggttgccg gtgtcagtgg ccaccgccga accgccccaa ggcgtgtcat cgggcacggt 1032
286 tggttctgtt tcttcaatga tctgtggcgc ttccaccttg tttgtcatgg cgtctttcgc 1092
288 tgccatgacg gcaaaccata acaggttaagc gatgccaccc cagcgcataa tatcgagcac 1152
290 gatcggcgcg gcattggaca aaagatcaac gcccaagggt ccggcgatga acaaaaagac 1212
292 gtcagaaatt aaacacacga gaagaaccgc aatgagtcct tcgcgcttaa ttcttgttt 1272
294 aatcaccagt acattctgcg gtccgatgga cagtaaaaga ctggcccca aaagcagacc 1332
296 tgtaatgaag atttccatga tcaccatcgt gacctatgga agtacttaag taaaatgatt 1392
298 ggttcttaac atggtttaat atagcttc atg aac ccc att caa ctg gac act 1444
299      Met Asn Pro Ile Gln Leu Asp Thr
300      220      225
302 ttg ctc tca atc att gat gaa ggc agc ttc gaa ggc gcc tcc tta gcc 1492
303 Leu Leu Ser Ile Ile Asp Glu Gly Ser Phe Glu Gly Ala Ser Leu Ala
304      230      235      240
306 ctt tcc att tcc ccc tcg gcg gtg agt cag cgc gtt aaa gct ctc gag 1540
307 Leu Ser Ile Ser Pro Ser Ala Val Ser Gln Arg Val Lys Ala Leu Glu
308      245      250      255
310 cat cac gtg ggt cga gtg ttg gta tcg cgc acc caa ccg gcc aaa gca 1588
311 His His Val Gly Arg Val Leu Val Ser Arg Thr Gln Pro Ala Lys Ala
312      260      265      270
314 acc gaa gcg ggt gaa gtc ctt gtg caa gca gcg cgg aaa atg gtg ttg 1636
315 Thr Glu Ala Gly Glu Val Leu Val Gln Ala Ala Arg Lys Met Val Leu

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## RAW SEQUENCE LISTING

DATE: 08/12/2002

PATENT APPLICATION: US/09/105,117I

TIME: 14:05:45

Input Set : A:\Sequencelisting\_FZJ 9910 PCT\_US.txt

Output Set: N:\CRF3\08122002\I105117I.raw

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316      275      280      285
318 ctg caa gca gaa act aaa gcg caa cta tct gga cgc ctt gct gaa atc 1684
319 Leu Gln Ala Glu Thr Lys Ala Gln Leu Ser Gly Arg Leu Ala Glu Ile
320 290      295      300      305
322 ccg tta acc atc gcc atc aac gca gat tcg cta tcc aca tgg ttt cct 1732
323 Pro Leu Thr Ile Ala Ile Asn Ala Asp Ser Leu Ser Thr Trp Phe Pro
324      310      315      320
326 ccc gtg ttc aac gag gta gct tct tgg ggt gga gca acg ctc acg ctg 1780
327 Pro Val Phe Asn Glu Val Ala Ser Trp Gly Gly Ala Thr Leu Thr Leu
328      325      330      335
330 cgc ttg gaa gat gaa gcg cac aca tta tcc ttg ctg cgg cgt gga gat 1828
331 Arg Leu Glu Asp Glu Ala His Thr Leu Ser Leu Leu Arg Arg Gly Asp
332      340      345      350
334 gtt tta gga gcg gta acc cgt gaa gct aat ccc gtg gcg gga tgt gaa 1876
335 Val Leu Gly Ala Val Thr Arg Glu Ala Asn Pro Val Ala Gly Cys Glu
336      355      360      365
338 gta gta gaa ctt gga acc atg cgc cac ttg gcc att gca acc ccc tca 1924
339 Val Val Glu Leu Gly Thr Met Arg His Leu Ala Ile Ala Thr Pro Ser
340 370      375      380      385
342 ttg cgg gat gcc tac atg gtt gat ggg aaa cta gat tgg gct gcg atg 1972
343 Leu Arg Asp Ala Tyr Met Val Asp Gly Lys Leu Asp Trp Ala Ala Met
344      390      395      400
346 ccc gtc tta cgc ttc ggt ccc aaa gat gtg ctt caa gac cgt gac ctg 2020
347 Pro Val Leu Arg Phe Gly Pro Lys Asp Val Leu Gln Asp Arg Asp Leu
348      405      410      415
350 gac ggg cgc gtc gat ggt cct gtg ggg cgc agg cgc gta tcc att gtc 2068
351 Asp Gly Arg Val Asp Gly Pro Val Gly Arg Arg Arg Val Ser Ile Val
352      420      425      430
354 ccg tcg gcg gaa ggt ttt ggt gag gca att cgc cga ggc ctt ggt tgg 2116
355 Pro Ser Ala Glu Gly Phe Gly Glu Ala Ile Arg Arg Gly Leu Gly Trp
356      435      440      445
358 gga ctt ctt ccc gaa acc caa gct gct ccc atg cta aaa gca gga gaa 2164
359 Gly Leu Leu Pro Glu Thr Gln Ala Ala Pro Met Leu Lys Ala Gly Glu
360 450      455      460      465
362 gtg atc ctc ctc gat gag ata ccc att gac aca ccg atg tat tgg caa 2212
363 Val Ile Leu Leu Asp Glu Ile Pro Ile Asp Thr Pro Met Tyr Trp Gln
364      470      475      480
366 cga tgg cgc ctg gaa tct aga tct cta gct aga ctc aca gac gcc gtc 2260
367 Arg Trp Arg Leu Glu Ser Arg Ser Leu Ala Arg Leu Thr Asp Ala Val
368      485      490      495
370 gtt gat gca gca atc gag gga ttg cgg cct tag ttacttctga aaaggttcag 2313
371 Val Asp Ala Ala Ile Glu Gly Leu Arg Pro
372      500      505
374 ggtttttcac ttcttcgcc gcaggaattg ggccaggcag agtaacacct tcagcaaatg 2373
376 g 2374

```

09/105,117I

4

<210> 5 → last sequence in submitted file  
 <211> 290  
 <212> PRT (LysG) either delete this or move it to C2207-C2237  
 <213> Corynebacterium glutamicum section

&lt;400&gt; 5

Met Asn Pro Ile Gln Leu Asp Thr Leu Leu Ser Ile Ile Asp Glu Gly  
 1 5 10 15

Ser Phe Glu Gly Ala Ser Leu Ala Leu Ser Ile Ser Pro Ser Ala Val  
 20 25 30

Ser Gln Arg Val Lys Ala Leu Glu His His Val Gly Arg Val Leu Val  
 35 40 45

Ser Arg Thr Gln Pro Ala Lys Ala Thr Glu Ala Gly Glu Val Leu Val  
 50 55 60

Gln Ala Ala Arg Lys Met Val Leu Leu Gln Ala Glu Thr Lys Ala Gln  
 65 70 75 80

Leu Ser Gly Arg Leu Ala Glu Ile Pro Leu Thr Ile Ala Ile Asn Ala  
 85 90 95

Asp Ser Leu Ser Thr Trp Phe Pro Pro Val Phe Asn Glu Val Ala Ser  
 100 105 110

Trp Gly Gly Ala Thr Leu Thr Leu Arg Leu Glu Asp Glu Ala His Thr  
 115 120 125

Leu Ser Leu Leu Arg Arg Gly Asp Val Leu Gly Ala Val Thr Arg Glu  
 130 135 140

Ala Asn Pro Val Ala Gly Cys Glu Val Val Glu Leu Gly Thr Met Arg  
 145 150 155 160

His Leu Ala Ile Ala Thr Pro Ser Leu Arg Asp Ala Tyr Met Val Asp  
 165 170 175

Gly Lys Leu Asp Trp Ala Ala Met Pro Val Leu Arg Phe Gly Pro Lys  
 180 185 190

Asp Val Leu Gln Asp Arg Asp Leu Asp Gly Arg Val Asp Gly Pro Val  
 195 200 205

Gly Arg Arg Arg Val Ser Ile Val Pro Ser Ala Glu Gly Phe Gly Glu  
 210 215 220

Ala Ile Arg Arg Gly Leu Gly Trp Gly Leu Leu Pro Glu Thr Gln Ala  
 225 230 235 240

Ala Pro Met Leu Lys Ala Gly Glu Val Ile Leu Leu Asp Glu Ile Pro  
 245 250 255

Ile Asp Thr Pro Met Tyr Trp Gln Arg Trp Arg Leu Glu Ser Arg Ser  
 260 265 270

Do not show  
 any

extraneous

responses to

C2127.

The only  
 correct  
 responses  
 are: DNA, RNA,  
 or PRT

(see 1.823 of  
 Sequence Rules)

Please  
 edit all  
 similar  
sequences

see  
 next page for  
 more errors

09/105,117I 5

Leu Ala Arg Leu Thr Asp Ala Val Val Asp Ala Ala Ile Glu Gly Leu  
275 280 285

Arg Pro  
290

FZJ 9910 PCT/US 6/8

FZJ 9910 PCT/US 8/8

*delete*

09/105, 117I

6

<210> 3  
<211> 2374  
<212> DNA (complement to <210> 1)  
<213> Corynebacterium glutamicum  
<220>  
<221> unsure  
<222> CDS (2) .. (652)  
<223> orf3  
<220>  
<221> gene  
<222> CDS (1421) .. (2293)  
<223> LysG  
  
<400> 3

either  
delete this or show it in <220> -  
<223>  
section

Otherwise, the sequence is  
considered  
erroneous

VERIFICATION SUMMARY

DATE: 08/12/2002

PATENT APPLICATION: US/09/105,117I

TIME: 14:05:46

Input Set : A:\Sequencelisting\_FZJ 9910 PCT\_US.txt

Output Set: N:\CRF3\08122002\I105117I.raw

L:10 M:270 C: Current Application Number differs, Replaced Current Application Number  
L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:206 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQ ID NO  
L:206 M:283 W: Missing Blank Line separator, <210> field identifier  
L:208 M:283 W: Missing Blank Line separator, <220> field identifier  
L:212 M:283 W: Missing Blank Line separator, <220> field identifier  
L:217 M:282 W: Numeric Field Identifier Missing, <212> is required.  
L:217 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQUENCE ID NOS:0 differs:3  
L:13 M:203 E: No. of Seq. differs, <160> Number Of Sequences:Input (2) Counted (5)